

SEQUENCE LISTING**SEQ ID NO: 1 (MOUSE GCR1/FRAGILIS NUCLEIC ACID)**

Mouse GCR1 (Fragilis) full length nucleotide sequence

5 GCCGCAGAAAGGGCAGACCCGACGCGCTCCATCCTTTGCCCTCCAGTGCTGCCTTTGCTCCGC
 ACCATGAACCACACTTCTCAAGCCTTCATCACCGCTGCCAGTGGAGGACAGCCCCAAACTACGA
 AAGAATCAAGGAAGAATATGAGGTGGCTGAGATGGGGGCACCGCACGGATCGGCTTCTGTCAGAA
 CTACTGTGATCAACATGCCCAGAGAGGTGTCGGTGCCTGACCATGTGGTCTGGTCCCTGTTCAAT
 ACACCTTTTCATGAACCTTCTGCTGCCTGGGCTTCATAGCCTATGCCTACTCCGTGAAGTCTAGGGA
 10 TCGGAAGATGGTGGGTGATGTGACTGGAGCCCAGGCCTACGCCTCCACTGCTAAGTGCTGAACA
 TCAGCACCTTGGTCCCTCAGCATCCTGATGGTTGTTATCACCATTGTTAGTGTGCATCATCATTGTT
 CTTAACGCTCAAAACCTTCACACTTAATAGAGGATTCCGACTTCCGGTCTGAAGTGCTTCACCC
 TCCGCAGCTGCGTCCCTCCTTGCCCTCCCTACACGCAGGTGTAACACTCATTATCTATCCACA
 GTGGATTCAATAAAGTGCACCTTGATAACCACC

SEQ ID NO: 2 (MOUSE GCR1/FRAGILIS AMINO ACID)

15 Mouse GCR1 (Fragilis) amino acid sequence

MNHTSQAFITAASGGQPPNYERIKEEYEVAEMGAPHGSASVRRTTVINMPREVSVDPDHVVWSLFNT
 LFMNFCCLGFIAYAYSVKSRDRKMVGDTVGAQAYASTAKCLNISTLVLSILMVVITIVSVIIIVL
 NAQNLHT

SEQ ID NO: 3 (MOUSE GCR2/STELLA NUCLEIC ACID)

20 Mouse GCR2 (Stella) full length nucleotide sequence

GGATCACAGACTGACTGCTAATTGGGTCCTTGGTTTTAGGTCTTTTCAAAGACTAAGCAATCTTGT
 TCCGAGCTAGCTTTTGAGGCTTCTGCCCATCGCATCGCCATGGAGGAACCATCAGAGAAAGTCGA
 CCCAATGAAGGACCCTGAAACTCCTCAGAAGAAAGATGAAGAGGACGCTTTGGATGATACAGACG
 25 TCCTACAACCAGAAACACTAGTAAAGGTCATGAAAAAGCTAACCCTAAACCCCGGTGTCAAGCGG
 TCCGCACGCCCGCGCAGTCTACGGAACCGCATTGCAGCCGTACCTGTGGAGAACAAGAGTGAAAA
 AATCCGGAGGGAAGTTCAAAGCGCCTTTCCCAAGAGAAGGGTCCGCACTTTGTTGTGCGGTGCTGA
 AAGACCCTATAGCAAAGATGAGAAGACTTGTTCGGATTGAGCAGAGACAAAAAGGCTCGAAGGA
 AATGAGTTTGAACGGGACAGTGAGCCATTGAGATGTCTCTGCACTTTCTGCCATTATCAAAGATG
 GGATCCCTCTGAGAATGCGAAAATCGGGAAGAATTAGGAGCTTACATTGTACGCTGCCCTGGCTG
 30 TCGACGATGCCGCACAGCAGATGTGAAAGCTATTTTTTGTTTAAGATTAACTTTTTCTGGTGCT
 GGGAAATCTTAACCTTGTTAACTTTAAATTGTAGATAGGATGCACAACGATCCAGATTTATGTGA
 AGTTTGAAGCCTCAAGCTGTGAGGCCCGAGGGCTGAGGAATAAAGTAAATAGAATTTGGAGTATG
 TACGTTCTAATTTCCAGAAATTTGTAATAAAAGCATTTTTTGT

SEQ ID NO: 4 (MOUSE GCR2/STELLA AMINO ACID)

35 Mouse GCR2 (Stella) amino acid sequence

MEEPSEKVDPMKDPETPQKKDEEDALDDTDVLQPETLVKVMKKLTLPNGVKRSARRRSLRNRIAA
VPVENKSEKIRREVQSAFPKRRVRTLLSVLKDPIAKMRRLLVRIEQRQKRLEGNEFERDSEPFRCCL
CTFCHYQRWDPSENAKIGKN

SEQ ID NO: 5 (Rat GCR2 HOMOLOGUE NUCLEIC ACID)

- 5 Rat GCR2 (Stella) homologue genomic sequence; similar intron-exon structure as mouse-Stella. AC094826 contig No.5 (22671 - 27595: contig of 4925 bp in length)

CCACCTCCGACGTATGATGGCTCCTAGACGCA
ACACGAAGCGGACTCCCCGCATCATTCACGTAGACCCGCCTTCTGCTTTCCTGTGCGGGTTTTG
10 GGAAGCCCGGCGCCCTCTCTTCTCACCTTGCTCCACTAGCACGCGGCTGTTTTCACTGAGCCCA
GCACTGGCTAAGTGGAGCACCAGGAGTTTCAGGCTATCCTTCAGAGGGCAAGGTGTAGTCCATGG
TGGGCTACAGGAGACCCTCTCTCTCCGTGAGTACAGAGAGGCCAAACCCAAGCCAGACAGGGGTGA
TGATTAGGAACATACCTTCGTGCGGGGAGAAAATACCGGTTTCATATAGGAATAAGAGGAACCAGGA
GGTAGTTAAGGCTGTGGTGTCTGGTTGCGGGGTTTTTGACTCTCAACAACCACGTTTCAGAACGTG
15 CTGAGTTTTTATGATGGTGTAGAATTTCCCTTATCAGCAATTGGTCTCCGCGGTGTTTTCTTTTTCT
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25 TCCATTCCTGGTAAATGACTCTGAGGGGAAATAGGAACCCAGAATAGATTGAGCCGGGGGCTACC
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TGATACCTCCTAGCACTCCGGGCTGAGGGCGTGGCTCGGGAGGAGCCATTCCTTTGGAGAGGAAA
ACAACCTGCTGGCCTTGAATCTGCCCTAATACCTGACAGTTACATGGGACCTCCTTATTTCCACAG
GATTCTTTAGTCTTTGTTTGGGAGATTTCAAATCTTGAGACTGCTCAACCCTTCCTGGCCTAAC
30 ACTCACAAGGCCAGGCTAGACCCAAATTCGTCAACCCCTTCTGTGTCCAAAACGGTGGGTGGCT
AGCTGGCTCACCCCTGGTGTCACTTTGCTTTAACAATTCGGAAGGTTGTGGTAAGTTTCCTGTAT
AAAATAGGACCATCTACTGGGTGTGGTCCCATGTAAAGCAAGGTTGGTTTCCCAAATACCCTGT
TTACATAGATGTCCGGAAGCATTGGAGCAGGTCAATTAGATTAGGTGGAACAGCCTGTTTTTG
GAAAGCTTTCCAGGGCGGAAATGAACCCAGAGGCATATTGGGCAAGCCCTCCGGCTTAAGCAAC
35 ACAATTGGCTGCAGGGGTCTCTGGAAGAGGTGTGAGACAAGAGAGAATATGCAGGTTTCAGGACC
TCTGAACCTAGAGTTAGGCTGCTGTAACATTGTAACATTGCTGTAAGCAGAACAGCCCATGGTAAG
AAGCTCAGTGGATCTCTACAAACACTAGGATATCTGCTCAGGGTTTATGACCAGGCCCTGTGCAT
ATGGTTTGTCTTCTGTTGGCCCCCTCTCTTGAAGAGGGGTGATTATCTGTTACCCACTTCCTTGTT
TCTCTGGGGTATTACCTTGCAAAATGCAAAATGATATACTTCACTAATGTCTCCATCTTCTGTTT
40 CAGAAATCCTACAACCAGAAACACTAGTAAAGGTCATGAAAAGCTAACCTGAACCCAGTGCC
AAGCCGACAAAATATCATCGTCGTCAAAGGGTTTCGTCTCCAGGTTAAGAGCCAGCCTGTGGAGAA
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TGTCCGTGCTGAAAGACCCCATAGCAAGGATGAGAAGATTGTTTCGGGTGAGTTGCGTTTGTGGG
CGGGGCATAGATCTAAGAGCAACTCTAGCCTCAGGAATGGCACCTAGGTTAAACAGGGAATGTAG
45 ACAAGGATAGTGAATACCTGTGATTCCCAGCTCAAGAAAACAAGCTCCAAGGCTATCCTCTACTG
CGCAGTCTGAAGCTGGCCAGAGCTATAGCAAAATGATAAGTCAGTATAACATTTATTTTTGGAT
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CGGTGCTAGGCTTTTGTAGTATAAGGCTGACTTAGACTGGATCTCAGAGCTGAAGTGGACCTGTT
AGTCTTTGTAGACCAGGCTGGGGTGGTTTCTGCTTTCTCAGCGCCTAGCTCACATAGTAGGCATT

TTAAC TTTGTCTTAATAGTAATTTGAGTAATTTTGT TTTTCTCTTGAAGATTGAGCAGAGACAAA
 GACAGCTTGAAGGAAATGAGGTAAATGCATATGGATGGGTAGGGTGTCTATGGATGGGTAGGGTGT
 TCTTGT TTTTACTGTTTCCTTAGACAAGGAGTGTGTATGTGGAGAGTTACCTTCTCAACACAGGG
 AATCTGGTTATTAAAGCAGTACTTTAAAAATAAATAAAATAAAATAAAATAAAAGCAGTAG
 5 AAGGGGATTACATTTCTTTGAGTTGCAATATCCTGATTAAACATTTTCTTTTCAGAGACGAGAT
 GAGCCATTGAGATGTCTCTGCAC TTTCTGCCATTATCAGAGATGGGATCCTTCTGAGAATGCTAA
 AATCGGGCAGAACCAGAAGAATTAGGGCAGTTTGAATTGTACACCGTCTTGGCGTTAACGGTG
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 10 TCTTCTAATTGCTAACCTTTAAATTATATAGGATGTGTGACATTTGGATTTCATGGGAATGACAGA
 TTTACCCAAGAATTGAGCATGAGTCAAAGCCTGGTAGTTTGATTTAGAAGGTAATTGGAATAAAT
 CTTTTTATTTTAGATTTTCTAGTTTGCAGAGAAATTTGTAATAAAGGCAAATTTGTTATCTTTAA
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 TGGGTGTGTGATGTTTCAAGATTCAATGTGTGCTACCCTGTATTTCTGCTTGAGGCAAGGTCTCCA
 TGAGGCCTAGCTGGTCTAACTCCTGGTCTGCCTTTTGT TTTTCCCCTGAGTTTGTACACCATAGG
 15 CTTGTGCGCAAGATCTGGAAGAGGCTTGATGTTTGTGTTTGTGCTGTGTAATAAACAATTGGTTG
 ACATATTCTAAAGTGTGGCACTGTATTGACCTGTCTGTCTCATGAGGAAGTTAATGACCGGAGC
 ATAATTGTATGCTTTATTTCTGAGAGAAGTGTGAGGAAAGGAGGAGTTAGGAAGAAAGCCCCAG
 GCTGGGGTTAAGAGCACTGGCTGCTTTTCCAGAGGTCCTGAGTTCAATTCCCAGCAATCACCTGG
 TGGCTCCCGAACATCTGTAAACAGGATCCAA TGCCCTCTTTTGGTGTGTCTAAGAACTCCCTAGGC
 20 ATGCAGAGGATTTTGT TTTTGT TTTTGT TTTTGT TTTTGT TTTTGT TTTTGT TTTTGT TTTTGT TTTTGT
 GGAACCGAACCCAGGGCCTTGCGCTTGCTAAGCAAGCGCTCTACCACTGAGCTAAATCCCCAAC
 CCTACAATGGCCTTTTCTACCTGCTTTTGAATTATCAATAAAAGACTGGGGCAAAGAAAGGCT
 GGAGTGAAATGAGAGAGAACATGTGAAGAGTAAATGAGAGAGAGCATGAGGGAATGAATGAGAGAG
 TGAATGTGAGAACGAATGTGAGAGCGAGTGAGAGAACATGAGAAGAACACGTTAAGAGTGAGTGA
 25 AGAGAGAATGTGAGGTGTGTATGAAGATTGTGTGTGGGGTTGGGGATTTAGCTCAGTGGTAGAGT
 GCTTGCCTAGGAAGCACAAGGCCCTGGGTTCCGTCCCCAGCTCCAAAAAAGACCCAAAAA
 AAAAAAAAAAAGATTGTGTGTGTGTGTGAAAGGAGAGTGCATGTGGTGTGTGTGAGATATGTG
 CAAGGTGTGTATCAAGAGTGTGTGTGAGAGTGAAAGGGTAATGAACAGAGGTGTGCATGAGCGTG
 GGAGTTTGAAGAAAGAAACAGCAATAAAAAAAAAAAGCAGAGTGCACGAGAGAATGCAGAGTGTG
 30 TGCAACCTCAAGCTGAGACAGAGACAGAGAGAAAGAGAGAGAGAGAGAGACTTTAAGCCTTGA
 AATTACCTGTGAGTTTGTACCCAAATAGTAGTCTGTGTATATTTATTTTGTAGCCTTCCAGATCCC
 TGCTTCCAGTGGAAGCTCTGATTCTATGTTGAGGCTGGACCCTGGCAATAGTGGGCTTCTTGAA
 AAATAGTCAAAGGAAACAGTGCTACACCATGGACTTAAGCCTTTAGACTCAGTTCTGGCTTCAAG
 AGCAGCTGTGAGAAAATAAGTGATGAACTACTTGCACTGCAACTCGAATC
 35

SEQ ID NO: 6 (Rat GCR2 HOMOLOGUE NUCLEIC ACID)

Rat GCR2 (Stella) homologue genomic sequence; different intron-exon structure
 from mouse-Stella (fused exons). AC097234 (131006 132449: contig of 1444 bp in
 length)

40 CCAGGATTGAGACGAGCTAGGCCTCATGCATGGAGACCTTGCTCAAGCAGAAATAAACAGGGTA
 GCACACATTGAACTCTGAACATCACGAGTGTGCACACACCCACACATGCATCTGTAAAAACGAG
 TCCCCATCTCCAATGGCTCGTTCTAATCTGTTCTGTGTATTTATTAAAGATAACAAATTTGCCTC
 TATTACAAATTTCTCTGCAAACTAGAAAAATCTAAAATAAAAGATCTATTCCAATTACCTTCTAAA
 TCAAACTACCGGGCTTTGACTCATGCTCAATTCTTGGGTAAATCTGTCAATCCCATGAATCCAAA
 45 TGTACACATCCTATATAATTTAAAGGTTAGCAAGTAGAGATTTCCCCAGCACCAGAAAGGTTT
 AATCTTAAACAAAAAACAGCTTTACATCTGCTGCATGGCACCCTTAACGGCAAGGACAGTGTA
 TGATTCAAACCTGCCCTAATTCTTCTGGTTCTGCCCAATTTTAGCATTCTCAGAAGGATCCCATCT
 CTGATAATGGCAGAAAGTACAGAGACATCTGAATGGCTCAACTCTTCTCTCATTTCTTCAAGCT
 GTCTTTGTCTCTGCTCAATCCGAACAAATCTTCTCATCTTGCTATGGGGTCTTTCAGCACCGAC

AACAGTGTGCGGACCCTTCTCTTGGGAAAGGCGCTTTGAACTCCCCTCATGATTCTTTCACTTCT
 GTTCTCCACAGGCTGGCTCTTAATCTGGAGACGAACCCTTTGACGAAGATGATATTTTGGCCGAT
 TGAGATAGAATATCAAAACAACATTTAACATTTAAATAACTTAACGATATACACACCTTTTTTTT
 TTCCACCTCCCCACACAGACAAAAACAACCCTATTTTTTCTTTACAACCCCGCCTAAGCAAGCG
 5 AAGCATTAGTAACCTGACCAATCATAGAAAGGAAACACCACCAGACCACATCAAATAAAATAAAAT
 CACCGCCCAACCCCAACCCCTATAAAAAACCCGCGGACCACACCACATATACTCCCCCCCCCGC
 ACCATCACTACATCACCTCTCCACCCATTCCCACCTCCCCCCCCAACATTAACCCACCCCATC
 ACGGAAACCCCAACACCAACAAATAAATTAGACACATCGCATTACATAAATTGACACAAGACCC
 ACCCCAAAAGAGCAGCAAGATTAGAGCCACATCCTCGGCCAACACAATACACTCAACCTGCAT
 10 AGTATCTATCTCCACCCCAACCTAGAAACAAAAATCTAATCAGCACCAGGCACCCAAGTATCAGC
 CACACTCAAAAACATACCCACCAATTAAACACGCCCCACCCACCAACAACCCACCCGCTGACA
 ACACACTTCGGAACCTACCCTCAACATCACCAAAAGCAATCGCAAGTTACGATGACTCCAACCACC
 TCACTCTCTCATTTG

SEQ ID NO: 7 (Rat GCR2 HOMOLOGUE NUCLEIC ACID)

15 Rat GCR2 (Stella) homologue genomic sequence; different intron-exon structure
 from mouse-Stella (fused exons). AC093991 (1 - 7657: contig of 7657 bp in length)

ACTGCAAGTAGTTCATCATTTACAGATCAAAAGAAAGAATAAAAAACAAGGTGTCATGATC
 CCTCCAAAAGAGTGGAACTTCAACTGCCAGATCCAAGATACTGAAATGGGTAGCATGCTGGAG
 AAAGAATTCAAAAGTTAGGTAGAGAATCTGGTTGAGCAGAGCACTTGCTTTCTTCCAGAGGATC
 20 TGAGTTCAAGTCCCAGGACCTATATCACAGTTTTCTGTAACCTTAGCTCCAGAGGGTCTGACACT
 TCTGTTCACTGTGGGCACCTGCATTACAGACAAACATAAAGTAGTTCATCACCTTTTTCACAGA
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 GGAGAGATGGTTCATCTGTTAGCCCATTTATTGCTCTTGAAGAGAACCCAGGGTCTCCATAGCA
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 25 CAGGCATACACAATGAACCTGCACACATACAAAAGTCCATAGAGCCATAGTTACCATTGTGAGCT
 CTGAGAACCAAATCCGTGTTCTCTGCAAGAGCGACATGCACGCTGAGAACCAGGCACCTTTCCCA
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 50 CTGGGGACCGAACCAGGGCTTGTGCTTGCTAGGCAAGCGCTCTACCACTGAGCTAAATCCCCA

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CAAATTTCTCTGCAAACTAGAAAATCTGAAAGATCTATTCCAATTACCTTCTAAATCAAACCTACC
40 AGGCTTTGACTCATGCTCAATTCCTGGGTAAATTTGTCATTTCGCATGAATCCAAATGTCACACAT
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55 AAAGCAATAGCCATGTAATAATGCCTAGATATAACTCTTCTTGTTCAGCAGCAAATGCATAAGC

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 15 AGATCTCATTACAAATGGCTGTGAGCCACTACGTAGTTGCTGGGAATTGAACTCAGGACCTCTGG
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 GCTGGGGATTGAACTCAGGACCTCTGGAAGAGCAGTCAGTGCTCTTAACCCCTTAGCCATCTCTC
 20 CAGTTCTAAAGGACAATGTTAATCGGGGCTGGCTCACAGGTTCAAGGTTTCACTCCATTATCAT
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 GATCCAAAGGCAGACCAAAAAAAGACTGGCTTACGGGCTTACCATAAGCAGCTAAGAGGAAGGT
 CTCAAAGCCCACCCTACAGTGGCATGTTCTCCAACAAGGCCACATCTCCTAATAGTGCCACTCCC
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 25 CCCCTGCTTTTACCTAAGCTCATTAGGCAGCAATATGCCTCTTATTGTTTGTAGCTCAGCATCCTG
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 ACACCAATGCCTAGAGAGATGCTCTTCTGTACATATCGCATGTGCAGAAGAAAGGGTGCCAGATC
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30 **SEQ ID NO: 8 (Rat GCR2 HOMOLOGUE NUCLEIC ACID)**

Rat GCR2 (Stella) homologue genomic sequence; different intron-exon structure from mouse-Stella (fused exons). AC103122 (11084 - 13244: contig of 2161 bp in length)

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SEQ ID NO: 9 (RAT GCR2 HOMOLOGUE NUCLEIC ACID)

Rat GCR2 (Stella) homologue genomic sequence; different intron-exon structure from mouse-Stella (fused exons). AC099436 (1 - 21688: contig of 21688 bp in length)

[illegible]

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